



OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/911,047

DATE: 02/27/2002
TIME: 10:55:08

Input Set : A:\E104720060-SeqLst-072301.txt
Output Set: N:\CRF3\02272002\I911047.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Erikson, Glen
4 Daksis, Jasmine
5 Picard, Pierre
7 <120> TITLE OF INVENTION: HOMOGENEOUS ASSAY OF BIOPOLYMER BINDING BY
8 MEANS OF MULTIPLE MEASUREMENTS UNDER VARIED CONDITIONS
10 <130> FILE REFERENCE: E1047/20060
C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/911,047
C--> 13 <141> CURRENT FILING DATE: 2001-07-23
15 <160> NUMBER OF SEQ ID NOS: 9
17 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

125 <210> SEQ ID NO: 9
126 <211> LENGTH: 15
127 <212> TYPE: DNA
128 <213> ORGANISM: Artificial Sequence
130 <220> FEATURE:
131 <223> OTHER INFORMATION: Description of Artificial Sequence: ssDNA probe wherein the
3' end of each base is covalently bonded to a lysine N-terminal leaving a free carboxyl group
133 <400> SEQUENCE: 9
E--> 134 tatagtagaa accac 15 ←

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/911,047

DATE: 02/27/2002
TIME: 10:55:09

Input Set : A:\E104720060-SeqLst-072301.txt
Output Set: N:\CRF3\02272002\I911047.raw

L:12 M:270 C: Current Application Number differs, Replaced Application Number
L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:134 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:15 SEQ:9



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/911,047

DATE: 02/27/2002

TIME: 10:55:46

Input Set : A:\PTO.AMC.txt
 Output Set: N:\CRF3\02272002\I911047.raw

3 <110> APPLICANT: Erikson, Glen
 4 Daksis, Jasmine
 5 Picard, Pierre
 7 <120> TITLE OF INVENTION: HOMOGENEOUS ASSAY OF BIOPOLYMER BINDING BY
 8 MEANS OF MULTIPLE MEASUREMENTS UNDER VARIED CONDITIONS
 10 <130> FILE REFERENCE: E1047/20060
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/911,047
 C--> 13 <141> CURRENT FILING DATE: 2001-07-23
 15 <160> NUMBER OF SEQ ID NOS: 9
 17 <170> SOFTWARE: PatentIn Ver. 2.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 50
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial Sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: Description of Artificial Sequence: derived from
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 34 <212> TYPE: DNA
 35 <213> ORGANISM: Artificial Sequence
 37 <220> FEATURE:
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 45 <210> SEQ ID NO: 3
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 47 <212> TYPE: DNA
 48 <213> ORGANISM: Artificial Sequence
 50 <220> FEATURE:
 51 <223> OTHER INFORMATION: Description of Artificial Sequence: derived from
 52 exon 10 of the human cystic fibrosis gene
 54 <400> SEQUENCE: 3
 55 tggcaccatt aaagaaaata tactcttgg tgtttcctat gatgaatata 50
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 15
 60 <212> TYPE: DNA
 61 <213> ORGANISM: Artificial Sequence
 63 <220> FEATURE:
 64 <223> OTHER INFORMATION: Description of Artificial Sequence: derived from

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PATENT APPLICATION: US/09/911,047

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Input Set : A:\PTO.AMC.txt
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65 exon 10 of the human cystic fibrosis gene
67 <400> SEQUENCE: 4
68 atatcatctt tggtg 15
72 <210> SEQ ID NO: 5
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74 <212> TYPE: DNA
75 <213> ORGANISM: Artificial Sequence
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78 <223> OTHER INFORMATION: Description of Artificial Sequence: derived from
79 exon 10 of the human cystic fibrosis gene
81 <400> SEQUENCE: 5
82 atatcatcta tggtg 15
86 <210> SEQ ID NO: 6
87 <211> LENGTH: 15
88 <212> TYPE: DNA
89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: Description of Artificial Sequence: derived from
93 exon 10 of the human cystic fibrosis gene
95 <400> SEQUENCE: 6
96 atatcggctt tggtg 15
100 <210> SEQ ID NO: 7
101 <211> LENGTH: 15
102 <212> TYPE: DNA
103 <213> ORGANISM: Artificial Sequence
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107 exon 10 of the human cystic fibrosis gene
109 <400> SEQUENCE: 7
110 ataccatatt tagtg 15
114 <210> SEQ ID NO: 8
115 <211> LENGTH: 15
116 <212> TYPE: DNA
117 <213> ORGANISM: Artificial Sequence
119 <220> FEATURE:
120 <223> OTHER INFORMATION: Description of Artificial Sequence: ssDNA probe wherein the
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121 base is covalently bonded to a lysine N-terminal leaving a free carboxyl group
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123 cacccaaagat gatat 15
126 <210> SEQ ID NO: 9
127 <211> LENGTH: 15
128 <212> TYPE: DNA
129 <213> ORGANISM: Artificial Sequence
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3' end of each
133 base is covalently bonded to a lysine N-terminal leaving a free carboxyl group
135 <400> SEQUENCE: 9
136 tatagtagaa accac 15

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/911,047

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Input Set : A:\PTO.AMC.txt

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L:12 M:270 C: Current Application Number differs, Replaced Application Number

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

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